

SEQUENCE LISTING

<110> JAPAN TOBACCO INC.

Syngenta Limited

<120> The rice restorer gene to the rice BT type cytoplasmic male sterility.

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<150> JP 2002-197560

<151> 2002-07-05

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<223> Oligonucleotide primer for amplification of G4003 HindIII marker sequence.

<400> 3

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<210> 4

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<223> Oligonucleotide primer for amplification of G4003 HindIII marker sequence.

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<223> Oligonucleotide primer for amplification of C1361 MwoI marker sequence.

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<223> Oligonucleotide primer for amplification of G2155 MwoI marker sequence.

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<223> Oligonucleotide primer for amplification of G291 MspI marker sequence.

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acatttttc ttccgaaact tccg 24

<210> 11

<211> 24

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<223> Oligonucleotide primer for amplification of R2303 BslI marker sequence.

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<210> 12

<211> 24

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<223> Oligonucleotide primer for amplification of R2303 BslI marker sequence.

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<223> Oligonucleotide primer for amplification of S10019 BstUI marker sequence.

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<223> Oligonucleotide primer for amplification of S10019 BstUI marker sequence.

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gtcacataga agcagatggg ttcc 24

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<223> Oligonucleotide primer for amplification of S10602 KpnI marker sequence.

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 <213> Oryza sativa L.
 <223> PCR marker G4003 HindIII
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<211> 2601

<212> DNA

<213> *Oryza sativa* L.

<223> PCR marker C1361 MwoI

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2601

<211> 1333

<212> DNA

<213> *Oryza sativa* L.

<223> PCR marker G2155 MwoI

<400> 21

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<212> DNA

<213> *Oryza sativa* L.

<223> PCR marker G291 MspI

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<212> DNA

<213> *Oryza sativa* L.

<223> PCR marker R2303 BslI

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<212> DNA

<213> Oryza sativa L.

<223> PCR marker BstUI

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<211> 676

<212> DNA

<213> Oryza sativa L.

<223> PCR marker S10602 KpnI

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<213> *Oryza sativa* L.

<223> PCR marker Tsp509I

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Leu	His	Met	Met	Ala	Asp	Asp	Arg	Gly	Gly	Gly	Ser	Pro	Pro	Asp
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Thr Met Val Lys Asn Gly Val Met Pro Asp Cys Met Thr Tyr Asn	260	265	270
Ser Ile Leu His Gly Tyr Cys Ser Ser Gly Gln Pro Lys Glu Ala	275	280	285
Ile Gly Phe Leu Lys Lys Met Arg Ser Asp Gly Val Glu Pro Asp	290	295	300
Val Val Thr Tyr Ser Leu Leu Met Asp Tyr Leu Cys Lys Asn Gly	305	310	315
Arg Cys Met Glu Ala Arg Lys Ile Phe Asp Ser Met Thr Lys Arg	320	325	330
Gly Leu Lys Pro Glu Ile Thr Thr Tyr Gly Thr Leu Leu Gln Gly	335	340	345
Tyr Ala Thr Lys Gly Ala Leu Val Glu Met His Gly Leu Leu Asp	350	355	360
Leu Met Val Arg Asn Gly Ile His Pro Asp His Tyr Val Phe Ser	365	370	375
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